

## Refilin holds the cap

Olivia Gay,<sup>1,2</sup> Fumihiko Nakamura<sup>3</sup> and Jacques Baudier<sup>1,2,\*</sup>

<sup>1</sup>INSERM Unité 1038; équipe Biomix; Commissariat à l'Energie Atomique; <sup>2</sup>Université Joseph Fourier; Grenoble, France; <sup>3</sup>Translational Medicine Division; Department of Medicine; Brigham and Women's Hospital; Harvard Medical School; Boston, MA USA

**T**he Reflins (RefilinA and RefilinB) are a novel family of short-lived actin regulatory proteins that are expressed during changes in cellular phenotype such as epithelial to mesenchymal transition (EMT). The Reflins promote the formation of actin- and myosin-rich perinuclear bundles that are characteristic of cellular phenotypic switches. In epithelial cells, RefilinB is upregulated in response to TGF $\beta$  stimulation and functions in organization of apical perinuclear actin fibers during early stage of the EMT process.<sup>1</sup> In fibroblasts, RefilinB stabilizes perinuclear parallel actin bundles which resemble actin cap.<sup>2</sup> Reflins bind and modulate the function of Filamin A (FLNA). Upon binding to Reflins, FLNA is capable of assembling actin filaments into parallel bundles, possibly by undergoing conformational changes at the C-terminal. Perinuclear actin structures determine nuclear shape, cell morphology, cell adhesion and possibly cell proliferation and gene regulation. Identifying the role of Reflins in organizing perinuclear actin networks provides additional insight in the process of intracellular mechanotransduction that regulate changes in cellular phenotype such as those observed during EMT.

### Refilin Proteins

The actin cytoskeleton is crucial for development as it controls cell division, membrane remodelling, cell migration and differentiation. These essential functions rely on the dynamic nature of the actin cytoskeleton. The mechanisms that determine actin cytoskeleton organization and

dynamics are controlled by a wide array of regulatory proteins. In this context, we have identified a new family of short-lived actin regulatory proteins, the Reflins, which are expressed during cell differentiation switches and serve as organizers of perinuclear actin networks. There are two known Reflin isoforms, ReflinA and ReflinB. ReflinA has a half-life of less than one hour and is transiently upregulated during differentiation of rat neural multipotent precursor cells into glial progenitors (unpublished data), while RefilinB is expressed during epithelial-mesenchymal transition (EMT) mediated by TGF $\beta$ .<sup>1</sup>

Rat ReflinA and ReflinB proteins display 40% identity and 48% similarity (Fig. 1). Reflins are capable of forming homodimers; the dimerization domain is located at the monomer's N-terminus.<sup>1</sup> The stability of Reflins is determined by their N-terminal sequences; mutation of the N-terminal domain alters the half-life of Reflins (see Fig. 1, unpublished data). Reflins exhibit greater stability in cells treated with the protease inhibitor MG132, suggesting that Reflins are subject to proteasomal degradation (unpublished data). Given the obvious impact of Reflins on actin dynamics, the mechanisms that determine the stability and degradation of Reflins deserve further investigation.

### Refilin Promotes FLNA-Dependent Actin Bundles

In cells, Reflins are stabilized upon interaction with flamins, which are actin-binding and cross-linking proteins. Vertebrate flamins are the only

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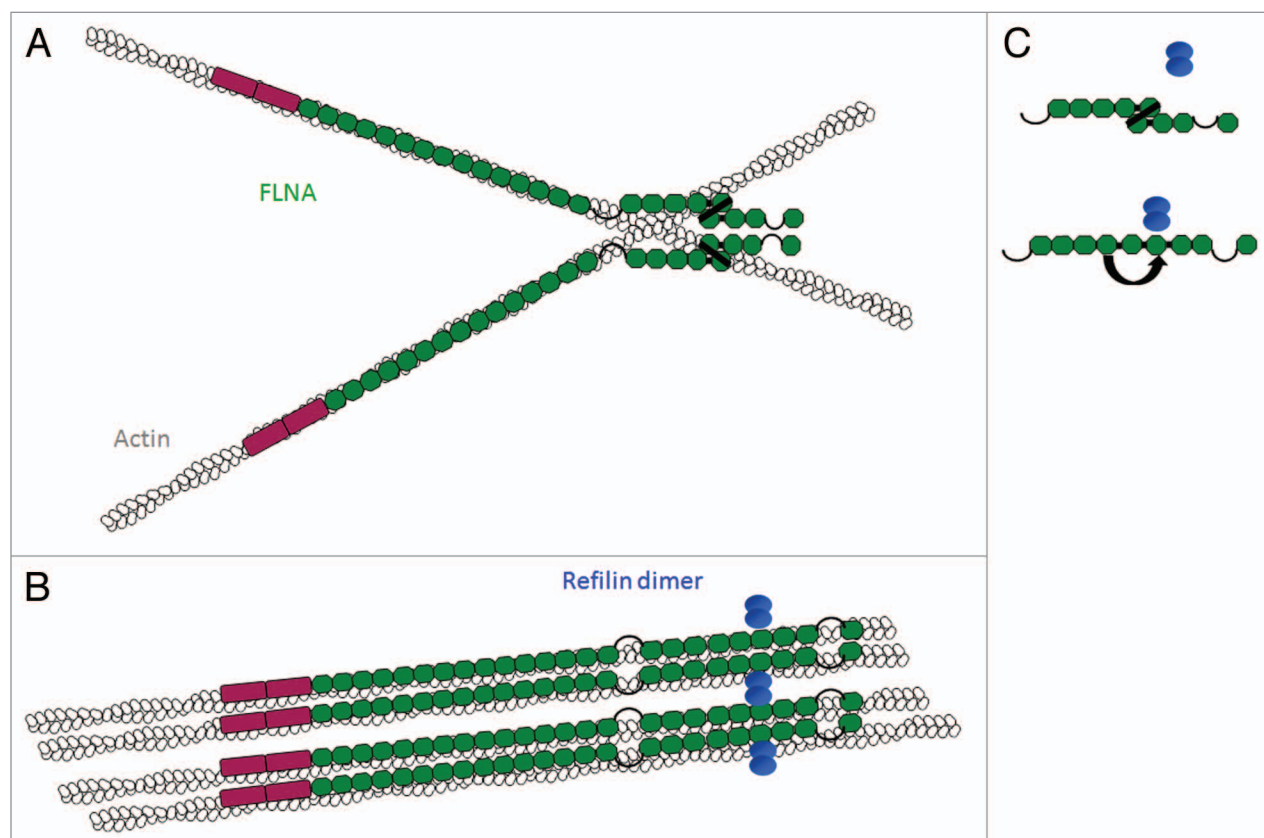
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\*Correspondence to: Jacques Baudier;  
Email: jacques.baudier@cea.fr

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ReflinA	MVGH	LHLQG	- - -	MGDTREQSR	DGLD	SPDSGLPPSPSPSP	PPFYAL	- - -	S	- - - - -	P	- -	G	45
ReflinB	MVGR	LSLQDVPEL	VDT	KKKG	-	DGVLDSPDSGLPPSPSPSH	- - WGLAAATGGGGERAPVAG	- - -	-	- - - - -	-	-	-	57
ReflinA	TLD	TRATTEVPAAT	SLFPNPPAL	- - - -	EMRS	- RLLPVFFGESIEVDPEPAHEIRCNSEV	- - -	-	-	- - - - -	-	-	-	99
ReflinB	TLE	PDAT	- - V	- - - T	SVVPNPASL	SHSLAGICSPRLCPLSFGEGVEFDPLPPKEIKYTSSV	- - -	-	-	- - - - -	-	-	-	112
ReflinA	TYASERYFRDKI	FYAPVP	- - -	TVTAYSETI	VAAPNCTWRSYRSQTL	EPRPRALRFGSTA	- - -	-	-	- - - - -	-	-	-	156
ReflinB	KYDSERHFI	DDV	- - -	QMPLGLVVASCSQTV	TCIPNCTWRNYKAEVRF	EPRHKPARFLSTT	- - -	-	-	- - - - -	-	-	-	169
ReflinA	IIFPKLARSSFR	TTL	- -	HC	- -	SLGQPRHWYSSSLQLRRCGDPAPSSGCPDVL	- - -	-	-	- - - - -	-	-	-	204
ReflinB	IIPKYPKT	VYTTTLDYNCHKKL	- - -	RR	-	FLSSVEL	- - -	-	-	EATEFLG	-	SDGLLDEC	-	216

**Figure 1.** Sequence alignment of rat ReflinA and ReflinB proteins. The two proteins show conserved regions with homologous (purple) or similar (blue) sequences. A 15 amino-acid N-terminal sequence is fully conserved between the two proteins, whereas a specific sequence is only found in ReflinB (red rectangle). These two regions function to control Reflin stability and degradation (manuscript in preparation).

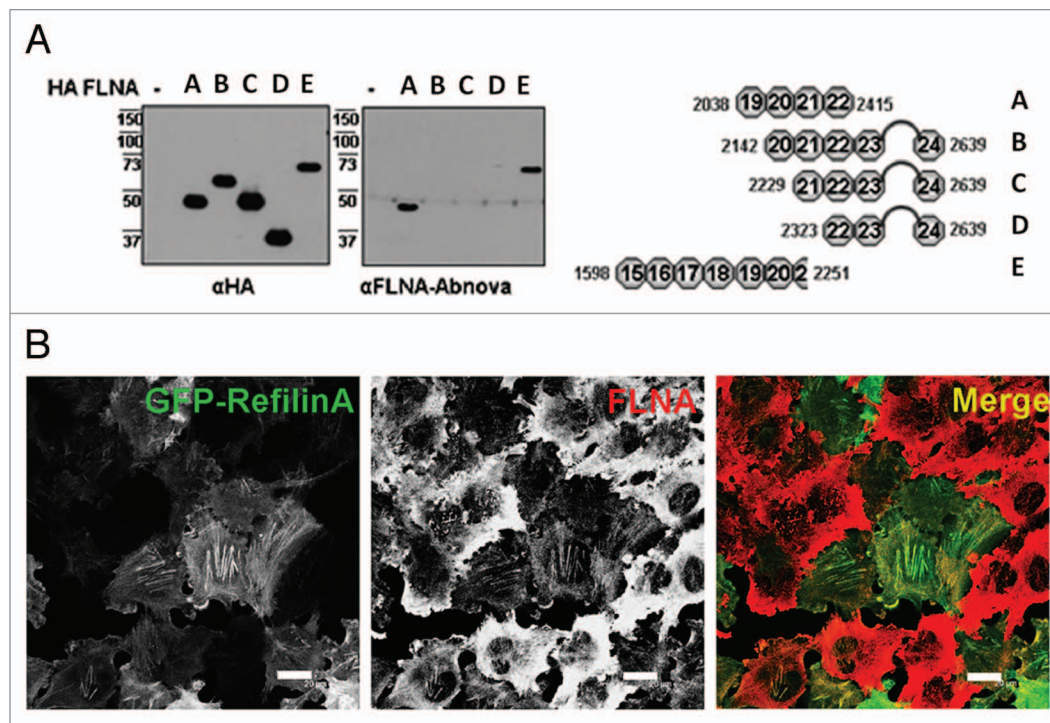


**Figure 2.** Reflin/FLNA complex organizes actin network into bundles. A proposed model by which Reflin binding promotes conformational changes in FLNA molecules that favor the bundling of actin filaments. (A) In the absence of bound Reflin, the V-shape FLNA molecule (green) generates actin networks (gray). (B) After binding of Reflin dimer (blue), FLNA acquires actin-bundling instead of actin-networking properties. (C) Domains 19, 20 and 21 on FLNA are folded such that domain 21 is entrapped between domains 19 and 20. The binding of dimeric Reflin to domain 21 of FLNA induces a conformational change of the 3 domains.

proteins known to co-immunoprecipitate with Reflins. Filamin A (FLNA) is the most abundant and best-characterized member of the three filamin isoforms. FLNA is ubiquitously expressed and provides mechanical stability to the actin cytoskeleton. FLNA also functions as a scaffolding protein for various cellular signaling pathways.<sup>3</sup> Vertebrate FLNA is

a homodimer of 280 kDa subunits composed of an N-terminal actin-binding domain followed by 24 immunoglobulin-like domains.<sup>4</sup> Two intervening calpain-sensitive “hinges” separate the repeats into rod 1 (repeats 1–15), rod 2 (repeats 16–23) and the dimerization domain (repeat 24). A secondary F-actin-binding domain resides in rod 1,<sup>4</sup> whereas rod 2

does not interact with F-actin, leaving it free to associate with partner proteins (Fig. 2). In the absence of bound Reflin, FLNA dimers crosslink actin filaments into orthogonal networks.<sup>4–7</sup> The binding of Reflins converts FLNA from an actin branching protein into one capable of assembling actin filaments into parallel bundles (Fig. 2A and B).



**Figure 3.** Monoclonal anti-FLNA antibody as a sensor of RefilinA-FLNA interaction. (A): Map depicting the epitopes of mouse monoclonal anti-FLNA. HA-tagged FLNA fragments (A–E) as indicated were expressed in human melanoma M2 cells and total cell lysates were probed with mouse anti-HA or anti-FLNA from Abnova. (B) U373 cells transfected with GFP-RefilinA plasmid were fixed and immunostained with an anti-FLNA antibody. Cells expressing GFP-RefilinA showed a drastic decrease in FLNA immunoreactivity. Bar = 20 μm.

Reflins enclose two putative FLNA binding domains and are capable of homodimerization at their N-termini.<sup>1</sup> It is therefore conceivable that Reflins cross-link FLNA and that crosslinked FLNA induces bundling. In this sense, ReflinA may function as a “zipper” that promotes the formation of multimolecular FLNA complexes on F-actin (Fig. 2B).

Moreover, three-dimensional structural studies revealed that the 19–21-domain fragment of human FLNA may function as an auto-inhibitory domain.<sup>8,9</sup> Based on detailed FLNA mutagenesis studies, we have proposed that Reflins bound to FLNA repeat 21 induce a conformational change in FLNA repeats 19–21 and change the high-angle F-actin branching mediated by FLNA into a low angle bundling of F-actin (Fig. 2). The notion that Reflin induces a conformational change in FLNA 19–21 domain is supported by the drastic decrease in FLNA immunoreactivity in the presence of ReflinA; this was determined by immunofluorescence using an antibody directed against repeat 19 of FLNA (Fig. 3).

### Reflin/FLNA Complex Localizes on Actomyosin Fibers

Reflin binding to FLNA promotes actin bundling in vitro.<sup>1</sup> In U373 astrocytoma cells, overexpression of a GFP-Refilin fusion protein promotes FLNA translocation from actin fibers, membrane ruffles and the cytoplasm to well-defined basal actin stress fibers that are connected to focal adhesion sites (Fig. 4). In addition, the Reflin/FLNA complex also appears on parallel perinuclear actomyosin fibers that organize above the nuclei and control nuclear height (Fig. 4B x-z and reviewed in ref. 1). In NIH 3T3 fibroblasts, endogenous ReflinB is expressed in confluent cells and the protein co-localizes with FLNA on perinuclear actomyosin fibers.<sup>1</sup> Co-localization of ReflinB with FLNA is also observed in basal stress fibers in overconfluent NIH 3T3 cells (unpublished data). In mouse epithelial NMuMG cells, ReflinB is expressed in response to TGFβ stimulation and the ReflinB/FLNA complex is specifically located on perinuclear actin filament bundles that form at the apical surface, but not on basal actin fibers.

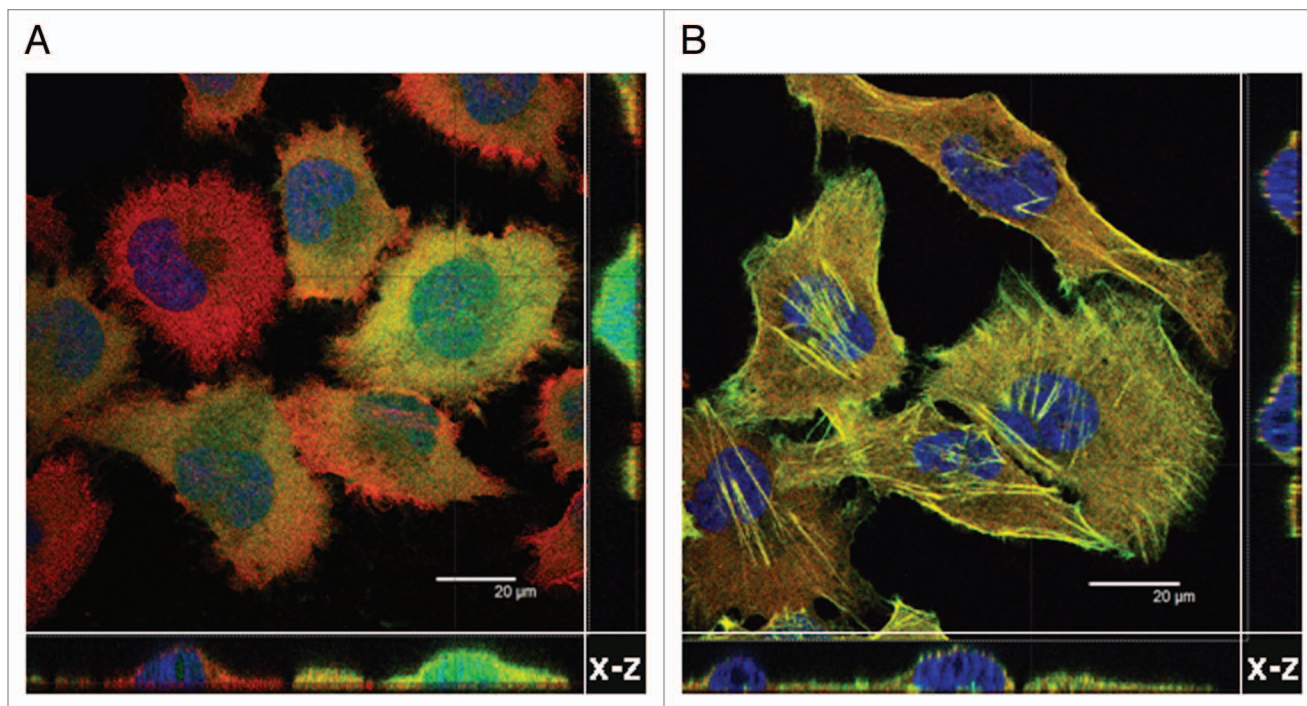
Collectively, these observations suggest that in the presence of low concentrations of ReflinB, the ReflinB/FLNA complex shows more avidity for the apical perinuclear actin cytoskeleton, although the ReflinB/FLNA protein complex may also localize to conventional basal actin fibers.

### What are Perinuclear Actin Structures?

Two perinuclear actin structures have recently been described: the “actin cap”<sup>2,10</sup> and the transmembrane actin-associated nuclear (TAN) line.<sup>11</sup> Actin caps generate tension in order to control nuclear shape while TAN lines are involved in nuclear positioning during cell polarization.

Both the actin cap and the TAN line are positive for myosin staining although they exhibit two main differences. First, the actin cap and TAN lines are oriented parallel and perpendicularly with the direction of cell migration, respectively. Second, actin caps are directly linked to focal adhesions; this is not observed in TAN lines. It is important to note that actin caps have been identified





**Figure 4.** Localization of the Reflin/FLNA complex in U373 MG cells. U373MG cells infected with recombinant adenovirus expressing GFP (A) or recombinant ReflinB-GFP (B) were fixed in paraformaldehyde and immunostained with antibodies against FLNA from USBiological (red). ReflinB forms a complex that can be found on basal stress fibers but also on apical perinuclear actin structures (x-z). Bar = 20  $\mu$ m.

in cells plated on micropatterned substrata<sup>2</sup> whereas TAN lines were observed in NIH3T3 fibroblasts migrating into scratch wounds.<sup>11</sup> Despite these differences, both actin caps and TAN lines-associated actin are linked to the nucleus by a specific set of proteins termed the LINC complex (linker of nucleoskeleton and cytoskeleton).<sup>12</sup>

In our studies with U373 and NIH 3T3 cells, the perinuclear actin structures associated with the Reflin/FLNA complex present characteristics of the actin cap. In epithelial NMuMG cells stimulated by TGF $\beta$ , the ReflinB/FLNA complex also contributes to the organization of apical perinuclear actin that accompanies the early stages of EMT. The identification of this novel perinuclear actin network provides additional insight into the mechanisms that regulate changes in cellular phenotype such as those observed during EMT.

## Conclusions

In eukaryotic cells, the actin perinuclear structures control nuclear movements and cell adhesion, which are essential functions for development. These structures may

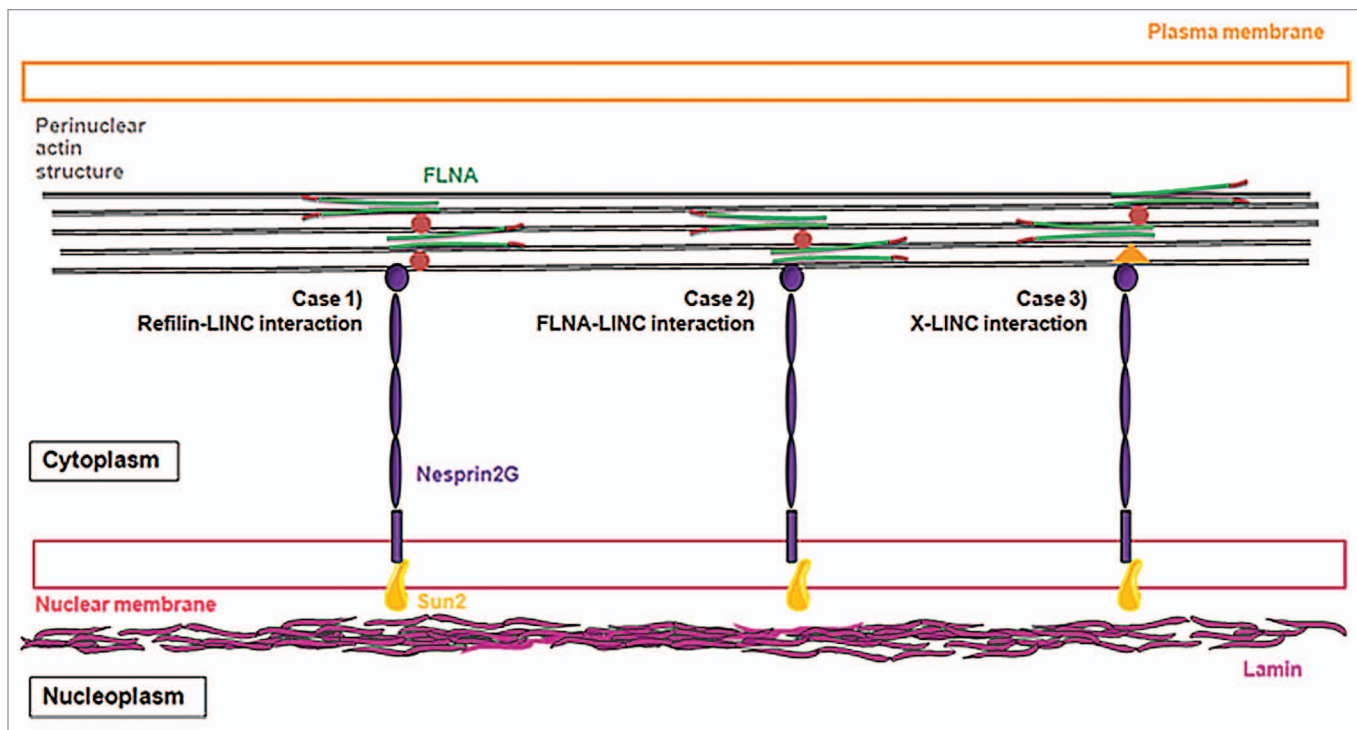
also influence gene expression and their presence correlates inversely with cellular proliferation.<sup>13,14</sup> The identification of an actin regulatory protein complex (Refilin/FLNA) that organizes perinuclear actin during changes in cellular phenotype has furthered our understanding of the role of perinuclear actin in normal and pathological situations. In Figure 5, we propose a model outlining the putative function of the Refilin/FLNA complex. The role of this protein complex in perinuclear actin organization requires further investigation.

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**Figure 5.** Hypothetical models of actin perinuclear structures stabilization by Refilin/FLNA complex. Schematic representation of the hypothesized roles of Refilin and FLNA in the formation of perinuclear actin structures. After binding to Refilin, FLNA changes from an actin-crosslinker to an actin bundler and the Refilin/FLNA complex subsequently organizes perinuclear actin structures by interacting with components of the LINC complex through binding to Refilin (blue), FLNA (green) or a presently unidentified protein (X, orange). These hypotheses will require further investigation.

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